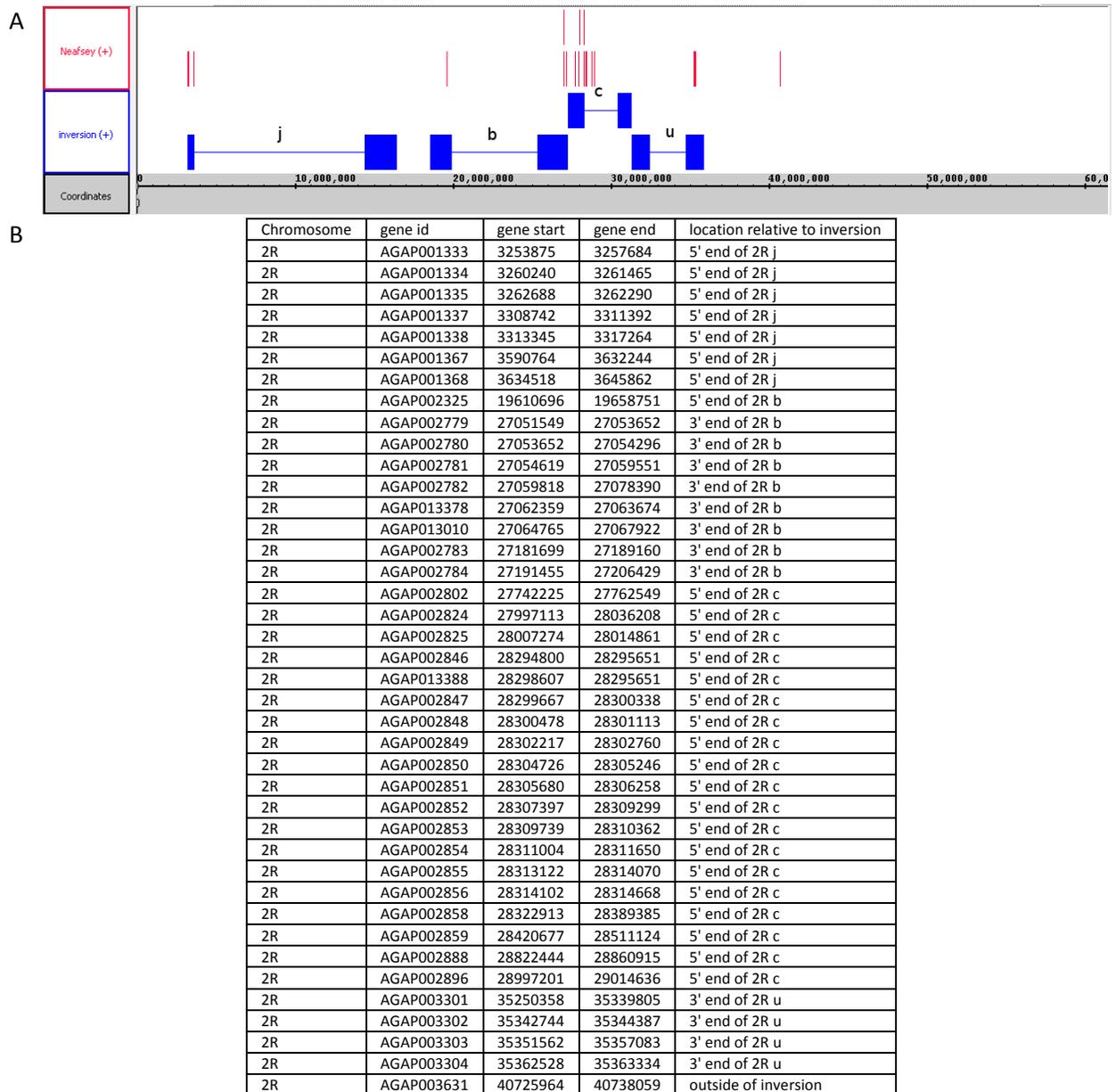


**Figure S3. A:** Genes reported to be diverged between *Bamako* and *Mali-S (Savanna)* form by Neafsey et al. [1] are marked in red. Inversion j, b, c, and u are marked with blue - blocks indicates uncertainty in starting and end of breakpoints. Most of the divergence was observed near breakpoints of inversions except for the 3' end of inversion c and 5' end of inversion u. **B:** Genes reported to be diverged between *Bamako* and *Savanna* form in the right arm of chromosome 2 by Neafsey et al. [1]



## REFERENCES

1. Neafsey DE, Lawniczak MK, Park DJ, Redmond SN, Coulibaly MB, et al. (2010) SNP genotyping defines complex gene-flow boundaries among African malaria vector mosquitoes. *Science* 330: 514-517.